

M C P S Q P T Y P G D P G P
ATG TGC CCG AGC CAG CCG ACC TAT CCG GGC GAT CCC GGG CCG
V E D L I R F Y D N L Q Q W
GTG GAA GAT CTG ATC CGC TTT TAT GAT AAC CTG CAG CAG TGG
L N C V T A A C *
CTG AAC TGC GTG ACC GCC GCC TGC TAG

FIGURE 1

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

1 11 21 31 41
ACACACCATA TGTGCCGAG CCAGCCGACC TATCCGGGCG ATCCCGGGCC
TGTGTGGTAT ACACGGGCTC GGTCGGCTGG ATAGGCCCGC TAGGGCCCGG

51 61 71 81 91
GGTGGAAAGAT CTGATCCGCT TTTATGATAA CCTGCAGCAG TGGCTGAAC
CCACCTTCTA GACTAGGCAG AAATACTATT GGACGTCGTC ACCGACTTGA

101 111 121 131
GCGTGACCGC CGCCTGCTAG GGATCCACAC AC
CGCACTGGCG CGGGACGATC CCTAGGTGTG TG

FIGURE 2

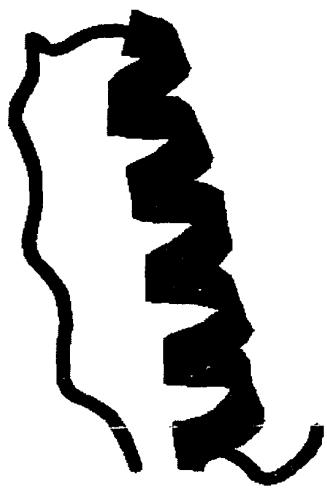


FIGURE 3

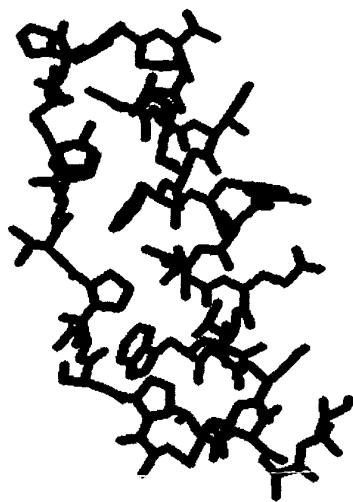


FIGURE 4

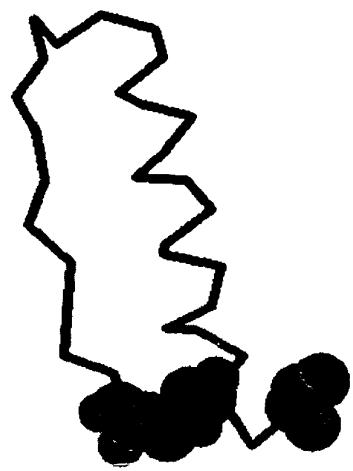


FIGURE 5

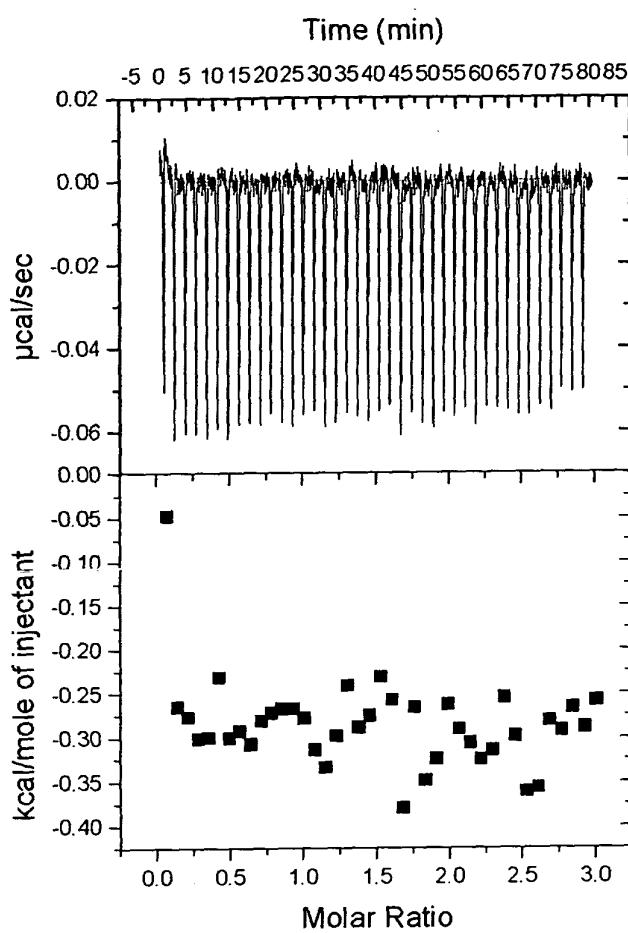


FIGURE 6

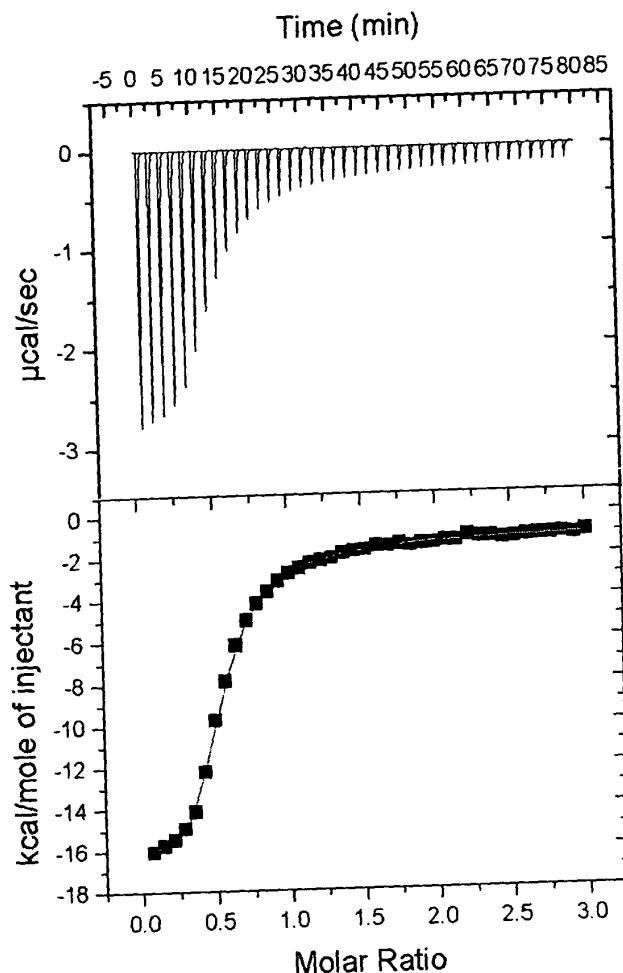


FIGURE 7

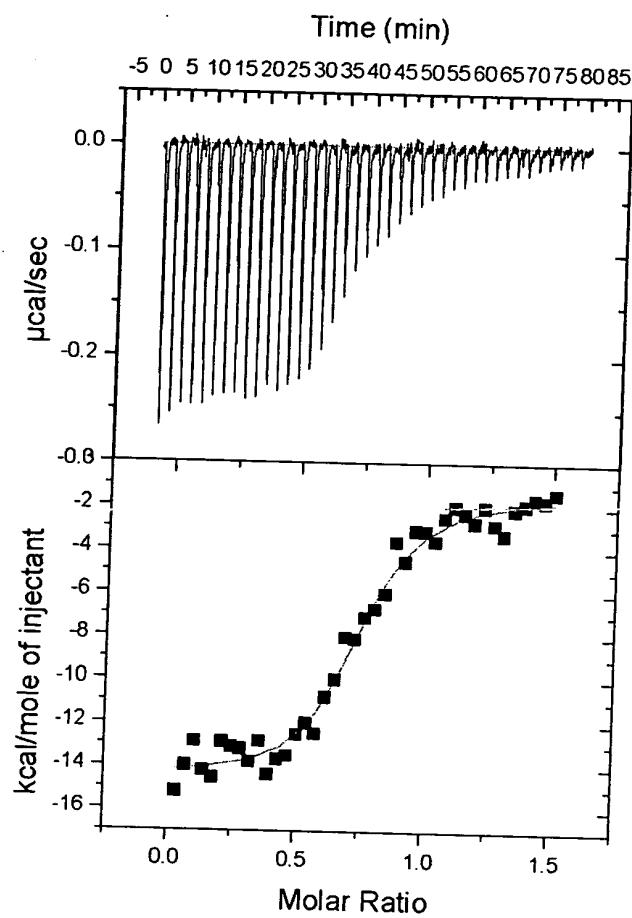


FIGURE 8

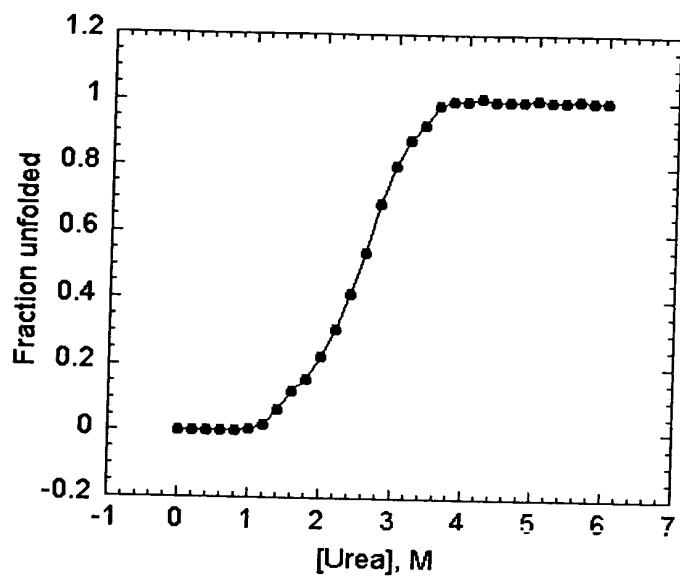


FIGURE 9

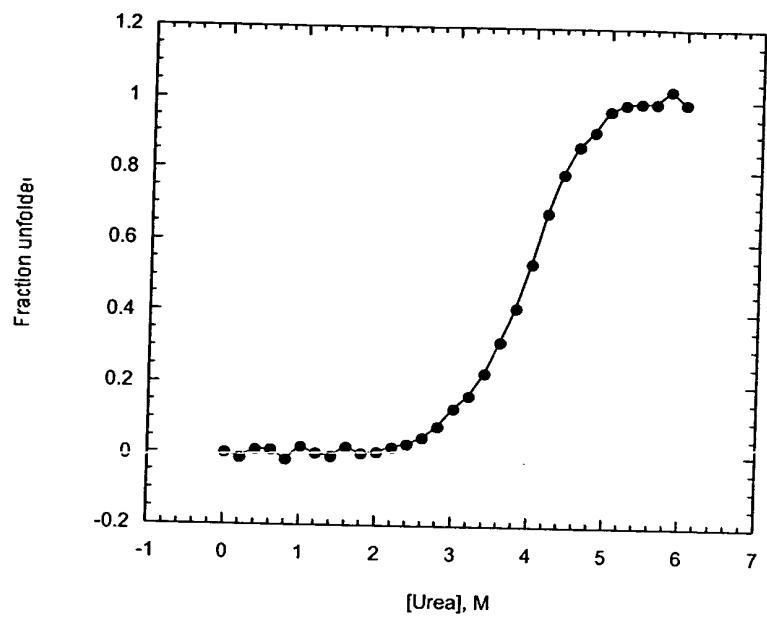


FIGURE 10

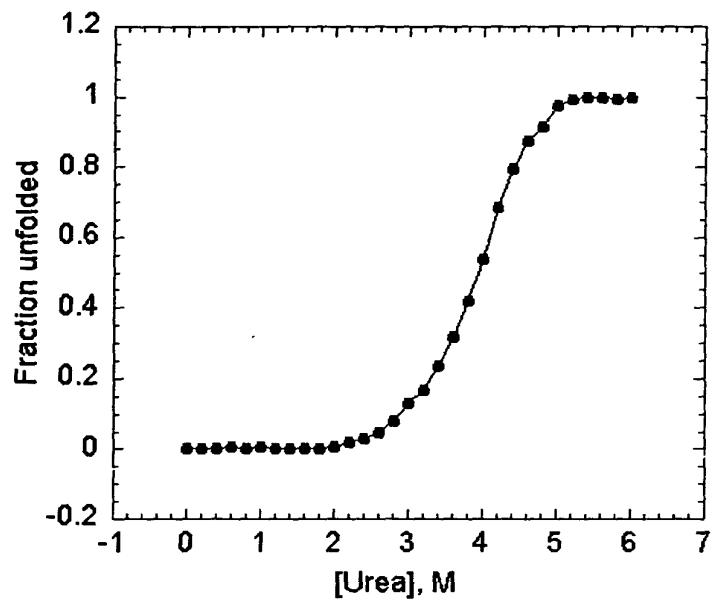


FIGURE 11

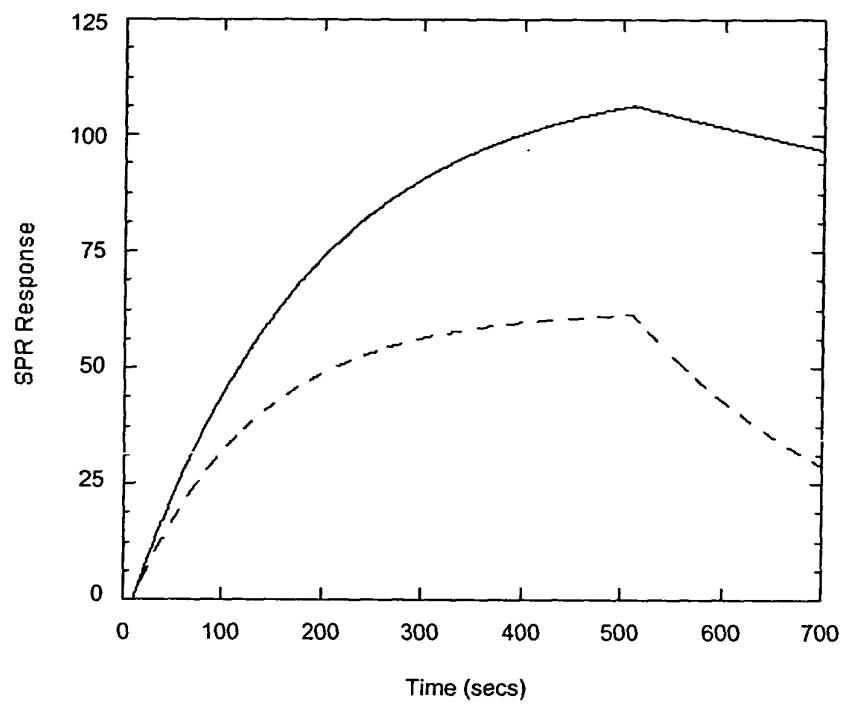


FIGURE 12

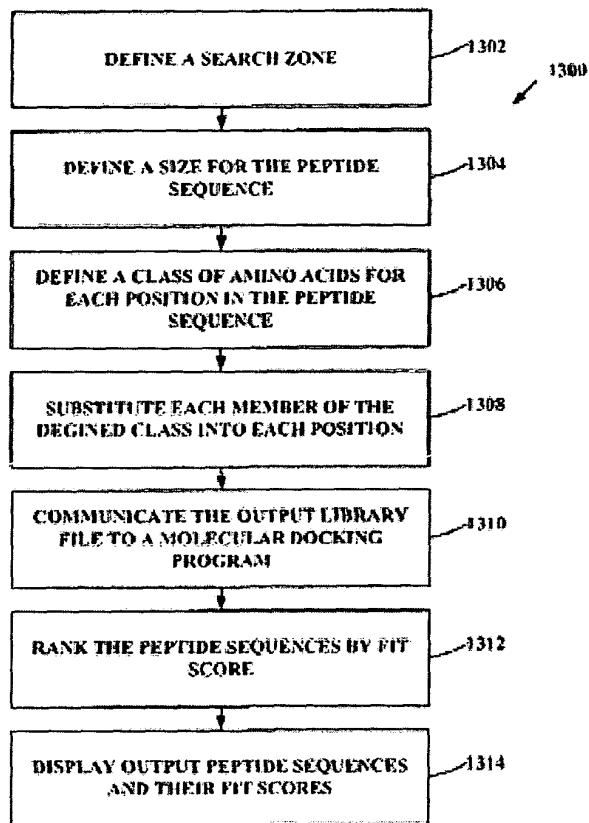


FIG. 13

4 000 02 70 393 - 4 12 2 6 0 34

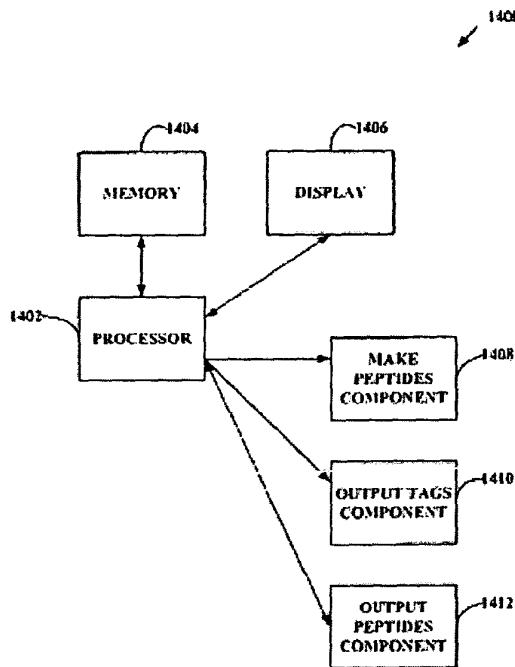


FIG. 14

4000270368 4226004

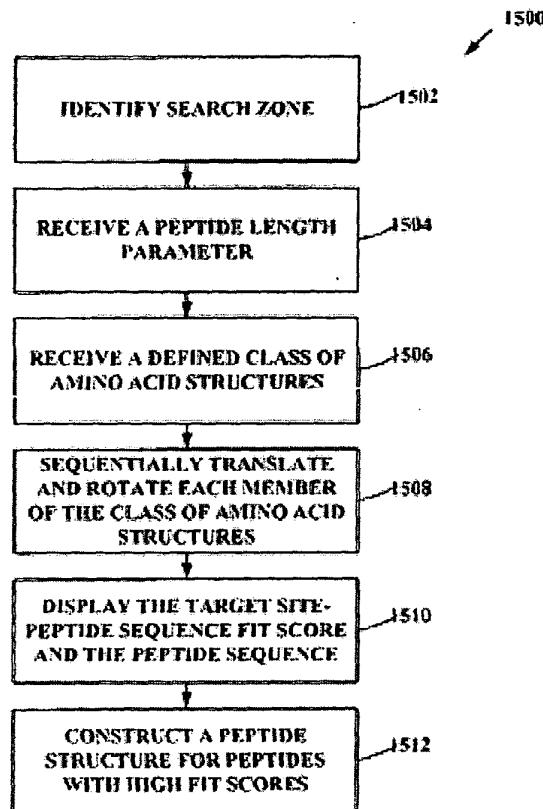


FIG. 15